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| <110> | Cooper, Richard K. Enright, Frederick M. Fioretti, William C. | |
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| <120> | Gene Therapy Using Transposon-Based Vectors | |
| <130> | 51687-0261 (331126) | |
| <140> <141> | US 10/583,812 2006-06-22 | |
| <150> <151> | PCT/US2004/43092 2004-12-24 | |
| | US 60/592,098 2004-07-28 | |
| | US 60/565,371 2004-04-26 | |
| | US 60/532,504 2003-12-24 | |
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6900
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agegeagaag tggteetgea actttateeg eetceateea gtetattaat tgttgeeggg
aagctagagt aagtagttcg ccagttaata gtttgcgcaa cgttgttgcc attgctacag
                                                                     7020
                                                                     7080
gcatcgtggt gtcacgctcg tcgtttggta tggcttcatt cagctccggt tcccaacgat
                                                                     7140
caaggcgagt tacatgatcc cccatgttgt gcaaaaaagc ggttagctcc ttcggtcctc
cgatcgttgt cagaagtaag ttggccgcag tgttatcact catggttatg gcagcactgc
                                                                    7200
                                                                    7260
ataattctct tactgtcatg ccatccgtaa gatgcttttc tgtgactggt gagtactcaa
                                                                    7320
ccaagtcatt ctgagaatag tgtatgcggc gaccgagttg ctcttgcccg gcgtcaatac
                                                                     7380
gggataatac cgcgccacat agcagaactt taaaagtgct catcattgga aaacgttctt
cggggcgaaa actctcaagg atcttaccgc tgttgagatc cagttcgatg taacccactc
                                                                     7440
                                                                    7500
gtgcacccaa ctgatcttca gcatctttta ctttcaccag cgtttctggg tgagcaaaaa
caggaaggca aaatgccgca aaaaagggaa taagggcgac acggaaatgt tgaatactca
                                                                    7560
                                                                    7620
tactcttcct ttttcaatat tattgaagca tttatcaggg ttattgtctc atgagcggat
acatatttga atgtatttag aaaaataaac aaataggggt tccgcgcaca tttccccgaa
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<400> 8 accatg

<211> 7

<212> DNA

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<220>

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<400> 9

| accatg | g | 7 |
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| <400> acgatg | 12 a | 7 |
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| <220> <223> | Kozak sequence | |

| <400> gacatga | 14 a | 7 |
|---------------------------|--|----|
| | 15 7 DNA Artificial Sequence | |
| <220> <223> | Kozak sequence | |
| <400> accatga | 15 a | 7 |
| <210><211><211><212><213> | 16 7 DNA Artificial Sequence | |
| <220> <223> | Kozak sequence | |
| <400> accatgt | 16 t | 7 |
| <211> <212> | | |
| <220> <223> | Base pairs 10651-11058 from GenBank Accession No Y00407 (Gallus sp.) | |
| <400> tctgcca | 17 attg ctgcttcctc tgcccttcct cgtcactctg aatgtggctt cttcgctact | 60 |
| gccacag | gcaa gaaataaaat ctcaacatct aaatgggttt cctgaggttt ttcaagagtc 1 | 20 |
| gttaag | caca tteetteece ageaceeett getgeaggee agtgeeagge accaacttgg 1 | 80 |
| ctactgo | ctgc ccatgagaga aatccagttc aatattttcc aaagcaaaat ggattacata 2 | 40 |
| tgcccta | agat cctgattaac aggcgtttgt attatctagt gctttcgctt cacccagatt 3 | 00 |
| atcccat | ttgc ctccc 3 | 15 |
| 210 | 1.0 | |

<211> 361

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      DNA
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<220>
<223> Synthetic
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                                                                      60
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ttggtttttt gtggatctgc tgtgccttct agttgccagc catctgttgt ttgcccctcc
                                                                     180
cccgtgcctt ccttgaccct ggaaggtgcc actcccactg tcctttccta ataaaatgag
                                                                     240
gaaattgcat cgcattgtct gagtaggtgt cattctattc tggggggtgg ggtggggcag
cacagcaagg gggaggattg ggaagacaat agcaggcatg ctgggggatgc ggtgggctct
                                                                     300
                                                                     360
atgggtacet etetetete etetetetet etetetetet eteteteteg gtacetetet
                                                                     361
C
<210> 19
<211>
      350
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<213> Artificial Sequence
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<223> Synthetic
<400> 19
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                                                                      60
ttatcatcac tttaaaaata aaaaacaatt actcagtgcc tgttataagc agcaattaat
                                                                     120
tatgattgat gcctacatca caacaaaaac tgatttaaca aatggttggt ctgccttaga
                                                                     180
                                                                     240
aagtatattt gaacattatc ttgattatat tattgataat aataaaaacc ttatccctat
                                                                     300
ccaagaagtg atgcctatca ttggttggaa tgaacttgaa aaaaattagc cttgaataca
                                                                     350
ttactggtaa ggtaaacgcc attgtcagca aattgatcca agagaaccaa
<210> 20
<211> 908
<212> DNA
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     Base pairs 1 - 1158 from GenBank Accession No. X00345 (Gallus
<223>
       sp.)
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<400>
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                                                                       60
                                                                      120
tgcatcagtt cagctacttg gctgcatttt gtatttggtt ctgtaggaaa tgcaaaaggt
                                                                      180
tctaggctga cctgcacttc tatccctctt gccttactgc tgagaatctc tgcaggtttt
                                                                      240
aattgttcac attttgctcc catttacttt ggaagataaa atatttacag aatgcttatg
aaacctttgt tcatttaaaa atattcctgg tcagcgtgac cggagctgaa agaacacatt
                                                                      300
                                                                      360
gatecegtga tttcaataaa tacatatgtt ceatatattg tttctcagta geetettaaa
                                                                      420
tcatgtgcgt tggtgcacat atgaatacat gaatagcaaa ggtttatctg gattacgctc
                                                                      480
tggcctgcag gaatggccat aaaccaaagc tgagggaaga gggagagtat agtcaatgta
gattatactg attgctgatt gggttattat cagctagata acaacttggg tcaggtgcca
                                                                      540
ggtcaacata acctgggcaa aaccagtctc atctgtggca ggaccatgta ccagcagcca
                                                                      600
gccgtgaccc aatctaggaa agcaagtagc acatcaattt taaatttatt gtaaatgccg
                                                                      660
                                                                      720
tagtagaagt gttttactgt gatacattga aacttctggt caatcagaaa aaggtttttt
                                                                      780
atcagagatg ccaaggtatt atttgatttt ctttattcgc cgtgaagaga atttatgatt
                                                                      840
gcaaaaagag gagtgtttac ataaactgat aaaaaacttg aggaattcag cagaaaacag
                                                                      900
ccacgtgttc ctgaacattc ttccataaaa gtctcaccat gcctggcaga gccctattca
                                                                      908
ccttcgct
<210>
       21
<211>
      901
<212>
      DNA
      Artificial Sequence
<213>
<220>
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sp.)
<400> 21
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ttctataact gaaatatatt tgctattgta tattatgatt gtccctcgaa ccatgaacac 120
tcctccagct gaatttcaca attcctctgt catctgccag gccattaagt tattcatgga 180
agatctttga ggaacactgc aagttcatat cataaacaca tttgaaattg agtattgttt 240

300

tgcattgtat ggagctatgt tttgctgtat cctcagaaaa aaagtttgtt ataaagcatt

Base pairs 431-1331 from GenBank Accession No. J00895 (Gallus

cacacccata aaaagataga tttaaatatt ccagctatag gaaagaaagt gcgtctgctc 360 ttcactctag tctcagttgg ctccttcaca tgcatgcttc tttatttctc ctattttgtc 420 480 aagaaaataa taggtcacgt cttgttctca cttatgtcct gcctagcatg gctcagatgc acgttgtaga tacaagaagg atcaaatgaa acagacttct ggtctgttac tacaaccata 540 600 gtaataagca cactaactaa taattgctaa ttatgttttc catctctaag gttcccacat 660 ttttctgttt tcttaaagat cccattatct ggttgtaact gaagctcaat ggaacatgag 720 caatatttcc cagtcttctc tcccatccaa cagtcctgat ggattagcag aacaggcaga 780 aaacacattg ttacccagaa ttaaaaacta atatttgctc tccattcaat ccaaaatgga 840 cctattgaaa ctaaaatcta acccaatccc attaaatgat ttctatggcg tcaaaggtca aacttctgaa gggaacctgt gggtgggtca caattcaggc tatatattcc ccagggctca 900 901 g

<210> 22

<211> 680

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic

<400> 22

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| gtgctgg | ggca gggcaatcca ttgccaccta tccc | aggtaa ccttccaact | gcaagaagat 6 | 560 |
|---------------------------|--|-------------------|----------------|-----|
| tgttgct | tac tctctctaga | | 6 | 580 |
| <210><211><212><212><213> | 23 72 DNA Artificial Sequence | | | |
| <220> <223> | Synthetic | | | |
| <400> gtggato | 23 caac atacagctag aaagctgtat tgcc | tttagc actcaagctc | aaaagacaac | 60 |
| tcagagt | tca cc | | | 72 |
| <210><211><211><212><213> | 24 62 DNA Artificial Sequence | | | |
| <220> <223> | From GenBank Accession No. J00 | 895 (Gallus sp.) | | |
| <400> | 24 | +a22aa +a2222aa | actoacact | 60 |
| | agct agaaagctgt attgccttta gcac | ccaage ccaaaagaca | acteagaget | |
| ca | | | | 62 |
| <210><211><212><212><213> | 25 1158 DNA Artificial Sequence | | | |
| <220> <223> | Base pairs 66 - 1223 from GenEsp.) | ank Accession No. | J00895 (Gallus | |
| <400> atgggct | 25 cca tcggcgcagc aagcatggaa tttt | gttttg atgtattcaa | ggagctcaaa | 60 |
| gtccac | catg ccaatgagaa catcttctac tgcc | ccattg ccatcatgtc | agctctagcc : | 120 |
| atggtat | cacc tgggtgcaaa agacagcacc agga | cacaga taaataaggt | tgttcgcttt : | 180 |
| gataaa | cttc caggattcgg agacagtatt gaag | ctcagt gtggcacatc | tgtaaacgtt 2 | 240 |
| cactctt | cas ttagagasat sstsaassaa atsa | ccaaac caaatgatgt | ttattcqttc 1 | 300 |

360 agccttgcca gtagacttta tgctgaagag agatacccaa tcctgccaga atacttgcag tgtgtgaagg aactgtatag aggaggcttg gaacctatca actttcaaac agctgcagat 420 caagccagag agctcatcaa ttcctgggta gaaagtcaga caaatggaat tatcagaaat 480 gtccttcagc caagctccgt ggattctcaa actgcaatgg ttctggttaa tgccattgtc 540 600 ttcaaaggac tgtgggagaa aacatttaag gatgaagaca cacaagcaat gcctttcaga 660 gtgactgagc aagaaagcaa acctgtgcag atgatgtacc agattggttt atttagagtg gcatcaatgg cttctgagaa aatgaagatc ctggagcttc catttgccag tgggacaatg 720 780 agcatgttgg tgctgttgcc tgatgaagtc tcaggccttg agcagcttga gagtataatc 840 aactttgaaa aactgactga atggaccagt tctaatgtta tggaagagag gaagatcaaa gtgtacttac ctcgcatgaa gatggaggaa aaatacaacc tcacatctgt cttaatggct 900 atgggcatta ctgacgtgtt tagctcttca gccaatctgt ctggcatctc ctcagcagag 960 1020 agcctgaaga tatctcaagc tgtccatgca gcacatgcag aaatcaatga agcaggcaga 1080 gaggtggtag ggtcagcaga ggctggagtg gatgctgcaa gcgtctctga agaatttagg 1140 gctgaccatc cattcctctt ctgtatcaag cacatcgcaa ccaacgccgt tctcttcttt 1158 ggcagatgtg tttcccct <210> 26 <211> 53 <212> DNA <213> Artificial Sequence <220> <223> Synthetic 53 <210> 27 <211> 103 <212> DNA <213> Artificial Sequence <220> <223> Synthetic <400> 27

60

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<211> 63
<212> DNA
<213> Artificial Sequence
<220>
<223> Base pairs 1145 - 1198 from GenBank Accession No. X00345 (Gallus
       sp.)
<400>
      28
                                                                      60
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                                                                      63
ggt
<210> 29
<211> 260
<212> DNA
<213> Artificial Sequence
<220>
<223> Base pairs 117 - 377 from GenBank Accession No. NM000207 (Homo
       sapiens)
<400> 29
                                                                      60
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                                                                     120
gaacgaggct tcttctacac acccaagacc cgccgggagg cagaggacct gcaggtgggg
                                                                     180
caggtggagc tgggcgggg ccctggtgca ggcagcctgc agcccttggc cctggagggg
tccctgcaga agcgtggcat tgtggaacaa tgctgtacca gcatctgctc cctctaccag
                                                                     240
                                                                     260
ctggagaact ctgcaactag
<210> 30
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic
<400> 30
Lys Tyr Lys Lys Ala Leu Lys Lys Leu Ala Lys Leu Leu
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<211> 39
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic
<400> 31
                                                                   39
aaatacaaaa aagcactgaa aaaactggca aaactgctg
<210> 32
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223>
     (Gly Pro Gly Gly) x where x is an integer from 3-9
<400> 32
Gly Pro Gly Gly
<210> 33
<211> 12
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic
<400> 33
Gly Pro Gly Gly Gly Pro Gly Gly Pro Gly Gly
               5
<210> 34
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic
<400> 34
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
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<211> 20
<212> PRT
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<223> Synthetic
<400> 35
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Gly Gly Gly Ser
<210> 36
<211> 5
<212> PRT
<213> Artificial Sequence
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<223> Synthetic
<400> 36
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